

1632

R. Shukla  
 RAW SEQUENCE LISTING DATE: 10/30/2000  
 PATENT APPLICATION: US/09/101,423A TIME: 11:56:31

Input Set : A:\Sequence\_Listing.txt  
 Output Set: N:\CRF3\10302000\x101423A.raw

**ENTERED**

3 <110> APPLICANT: Rudland, Philip S.  
 4 Barracough, Roger B.  
 W--> 5 <120> TITLE OF INVENTION: Metastasis Inducing DNA's  
 W--> 6 <130> FILE REFERENCE: WPT 0114 PUS  
 W--> 7 <140> CURRENT APPLICATION NUMBER: US 09/101,423A  
 8 <141> CURRENT FILING DATE: 1998-11-27  
 9 <150> PRIOR APPLICATION NUMBER: PCT/GB97/00074  
 10 <151> PRIOR FILING DATE: 1997-01-10  
 W--> 11 <160> NUMBER OF SEQ ID: 6  
 13 <210> SEQ ID NO: 1  
 14 <211> LENGTH: 1033 base pairs  
 15 <212> TYPE: DNA  
 16 <213> ORGANISM: Homo sapiens  
 18 <400> SEQUENCE: 1  
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 21 ctggacagac tctgggacag tccctctgc tctctgttg gcgcctgagt cccttttgc 120  
 23 ctggaggcccc ttacgttagc ctcccatctg gatgacact tagaaagct gggaaagggt 180  
 25 cacactcagg taactgagca gagtcaggat attaaaggat agtctggggg gcctcgagga 240  
 27 ttgatctgtc gccttaaaaa gccaatttgg tgactaaaccc agactattgt cactttaggt 300  
 29 gggaaaggcac tagcatatct gatgggtcac atctgagaad ggtttcttagc agtgggtggcc 360  
 31 ttgtgtgagc agcatggcgt gtatcatggt gtgcagcata ctcaggctgc ttgcaacact 420  
 33 cgaggctctt ttccatgtt aggggaacca ctgtgttgca acatgggtcca agaatacagt 480  
 35 catgtgagga gaatcccaat ggctcaggag aaaaacggagat tctgtgacct ccattcttca 540  
 37 agatacagaa ttattcttgg actgtgtttt catgctctt gtggatgggg gtgagttac 600  
 39 ttccatgttgc ttagcatgtc ttacttgtt tattcaagta aatgcttaaa ttatcttgg 660  
 41 tataccctctg tggaaacgac gtttttata catgcgcgtt gtccctgtga ttgataactgc 720  
 43 ttgaactcaa gagaactttt ctcatgtat ctttttttcaac cgtatggggat gaaactgtct 780  
 45 gatgtctca ataaagtgg ctctgtcagg agacgttagt ctgtccctgtt tatctgttcc 840  
 47 attcttccgc tcccaacggcc totacagcac taaacccacc accgatagac tcagtccttc 900  
 49 actgacaacaa atcaccagag gctcttaact gagattataa actgttaacta gatgtatgggt 960  
 51 tgaatcgctc cccagaaaca taaacatttta cttggagaac tcaagacccc ttttagaca 1020  
 53 taactcccat ggt 1033  
 55 <210> SEQ ID NO: 2  
 56 <211> LENGTH: 1058 base pairs  
 57 <212> TYPE: DNA  
 58 <213> ORGANISM: Homo sapiens  
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 60 attgtctgtga gcttatttgc gacatgggtt gacgcctt ttaagggggt agatacaaag 60  
 62 aatgggttga aatttctgtc cacaacgcgt ctccatgttt tcacaattac acttgcaacc 120  
 64 tgggtctcgc agccagaatt tagggatgtt atgggacagg gtcggggaaa gaaggagaag 180  
 66 ggttaaggaa agacagcagc ttaaagtccca aacagctccca ggagactate ttttagaaata 240  
 68 acatcagacc atgaggagaa ttgatatcat tttttttcaat tgggtatcgc caagggaaact 300  
 70 ttccatctgtt ttaaaaataaa ttactgttgg cactaatcc aattggaaat gccccacaca 360  
 72 atttatcttc cacttcatgtc tgcattccata tgcctgacgt ggcggagcag aagcatcc 420  
 74 tcccttctgtt ataaatgtt ttttgtaaat atttggagac gggatctgtt gtcacaggaa 480  
 76 acacgtacaa accggctgtt ttatctgtt cccgatagag gcccttcttg acgtacaggaa 540  
 78 ccccaaaaca gtcaggatgc ttttttttttccatgaa gcttggatc caatttagcaa 600

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|     |   |      |
|-----|---|------|
| 80  | ccattggagg aacgaggctg cactgtctac cacaaggcgc actttccaaa gagcacat       | 660  |
| 82  | atattggagc aagacatttt gctggctgac tggctgtg taagctgata aactgctata       | 720  |
| 84  | tttattaaac tggctttct ttgaacaccc cactcaagga aaaaaaaaca cacttaggt       | 780  |
| 86  | gacattttt ggatgtaag tctttataga gatgcttaag tttaaacagg acttttaaag       | 840  |
| 88  | ccggctctat tccatataat gaatgggtc octacaagg aagaaactgg gagcaggta        | 900  |
| 90  | tgtacactt gttgtgtgt agagacaac tgaggagtc aagaggagca cgtacaatgc         | 960  |
| 92  | agagaaggc tgaccctt tcacactgaa caaacatgc atgtgtgggt cgatagatga         | 1020 |
| 94  | gagtatcccc caaactcactc acatcgaaac gcttggtc                            | 1058 |
| 96  | <210> SEQ ID NO: 3  |      |
| 97  | <211> LENGTH: 1008 base pairs   |      |
| 98  | <212> TYPE: DNA   |      |
| 99  | <213> ORGANISM: Homo sapiens  |      |
| 101 | <400> SEQUENCE: 3   |      |
| 102 | aggaccagat ttacacatccc atcaaattggc ccagaagggtt ttaatgctgt cttttggccc  | 120  |
| 104 | aggggcgaac tgcacacaca tggcacata cactacaga gacacacatt cagcagcata       | 180  |
| 106 | agaacacaat cacaataaa aaaaattctt gaaaattttt agctaaaattt gttaaagaat     | 240  |
| 108 | aacatataat caattttctt ttatctttt aaaaattttt ttatataatg tatatgagta      | 300  |
| 110 | cactgcctct ccctccacatc atacgtatc agggcatcg atccattac agatgttgt        | 360  |
| 112 | gagccacat gtgtttcac agatgggtt gaggccatct gtggtttcg gaattgact          | 420  |
| 114 | caggacatcc tggaaagacg tcaatgtctt taacctctaa gccatcttc ctgaccctta      | 480  |
| 116 | tatacaattt taatgtctacg tacacacaaat tttctttcc ttaatgtttt gaggattttg    | 540  |
| 118 | tctggagaag taagaataaa gaggggaaag aacatgtctt tcacattgca ccagtggaa      | 600  |
| 120 | cagctgtttt aaatggtagaa tgccatggaa tgactggct gccttctat tactgttct       | 660  |
| 122 | caccatcttc cttttaactg gagctccctt atctaattta ttatgttgc gataccagg       | 720  |
| 124 | gtttctctt gtttgatct ttttaagaca gagactcacc atataccctt ggctggctg        | 780  |
| 126 | aagctcaacta tggatgaccat tctggccctt aactctaaagg agatctatctt gtttccatgt | 840  |
| 128 | gtctggatggaa aaggcttgcg ctacaaatgc tggctgtggg ctttggagca gcctcggtt    | 900  |
| 130 | tggctcttta taaggatctc taqcttagca gtaagttagcc tagccatgtt gttgttagaa    | 960  |
| 132 | gttggtcgtt catctggctt ccagcacaaa ggcagtcact aaacgtggc ctcatttcat      | 1008 |
| 134 | caagactgaa tgccaaattcc ttgtgtctt cctgtgtctt octggaaac                 |      |
| 136 | <210> SEQ ID NO: 4  |      |
| 137 | <211> LENGTH: 1088 base pairs   |      |
| 138 | <212> TYPE: DNA   |      |
| 139 | <213> ORGANISM: Homo sapiens  |      |
| 141 | <400> SEQUENCE: 4   |      |
| 142 | agtggggac acagctgtgt tgattaagat gtttcttggg aaaaggagtt aagectaatg      | 60   |
| 144 | atttccaataat gaaaggactg ctaattgggg aggcaatgtt gcttaatgg gacacctgcg    | 120  |
| 146 | ggttaattaa agctcttc cagtgccctt tccctttttt ggctctggg ggcgaaggca        | 180  |
| 148 | ttgaaqaggga tgcaggcatt ctaagggtt gtttcttggg tctcccttc cctctgtcc       | 240  |
| 150 | aactcagtga ggtatccctg tctgtgtgt cttcttagatg cctgtctgg gctctgtga       | 300  |
| 152 | gttaaggctt ctggatctgat gtcgtccatg gaaacacgtt gagtcattt gaaaggggag     | 360  |
| 154 | aaccaggcaaa aggtgttgcg tggatcttgc gaaattcttgg gggcaaaatgt tcaaggctaa  | 420  |
| 156 | ctctccattt agaccaatgt tgagactggc ctggaaacaa aaatataaaat tgagtggat     | 480  |
| 158 | cataatgacag cacctggatg tccctgtcc tagagatcat aaggacatgg ctgtgggg       | 540  |
| 160 | cttggtcgag atggactttt gttgtcgagg agggacatgg ccccaatgc ggaggccctg      | 600  |
| 162 | gaatgtccatc tggatataact gtggaaactg atgtgtctt tataatggaa gttgtgtgt     | 660  |
| 164 | tatctgtgtt acatctgtg tggtaatgtt tcacatgggtt ttatcttgcg gaggatattt     | 720  |
| 166 | cctggggaga aaaaaggact ggtggccacaa agcacaatgtt taaaatctgtt ggggtgtgac  | 780  |
| 168 | ccattatggaa ctatcataac tgagtgcagg tatcaagaat acttttagcag gtggtaaaaa   | 840  |

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|     |             |             |              |             |            |             |      |
|-----|-------------|-------------|--------------|-------------|------------|-------------|------|
| 170 | gatttttgaa  | tgcgcaacga  | ccaaaactga   | actaaaaat   | caagcatggc | atggatcctg  | 900  |
| 172 | gggtctcccg  | gaaggacttg  | cctttaactgc  | attgtgcgac  | ttgacggtag | ccttggttct  | 960  |
| 174 | gaatgcacaa  | cacgtggct   | ttgggtgcga   | caggccacca  | cgccgtgcct | gaaacacctc  | 1020 |
| 176 | agtcaggtt   | tgtggctatg  | tcctatgact   | tggacttact  | tttattgcac | atataaatat  | 1080 |
| 178 | tttccctgc   |             |              |             |            |             | 1088 |
| 180 | <210>       | SEQ ID NO:  | 5            |             |            |             |      |
| 181 | <211>       | LENGTH:     | 960          | base pairs  |            |             |      |
| 182 | <212>       | TYPE:       | DNA          |             |            |             |      |
| 183 | <213>       | ORGANISM:   | Homo sapiens |             |            |             |      |
| 185 | <400>       | SEQUENCE:   | 5            |             |            |             |      |
| 186 | gaggggggtgg | tggcacagtt  | atgtttttgt   | aggaagggtt  | ccatgaacct | cagcagagct  | 60   |
| 188 | cgggttagaa  | attttaaaacg | cctgaggggaa  | attttttttt  | taatcgtta  | tgaatctgac  | 120  |
| 190 | atgagaaaaa  | cagatcaga   | acgttcttgt   | gtttcagaaa  | aggacaagtg | tgtgagctaa  | 180  |
| 192 | cagactgcac  | actgggtttc  | gaggcacate   | tggatcacag  | gagcgtcaga | taatgtcccc  | 240  |
| 194 | aaagqtaat   | gcatttgctt  | gcacagtacc   | gagtgtgttgg | gggggtgcct | acagcccage  | 300  |
| 196 | ggttctcaac  | tttccctgat  | cttcgaccct   | ttaataactgt | gatttqata  | ctgttatgaa  | 360  |
| 198 | cccccaacott | aaaatattt   | ttgttgctgt   | tcataactgt  | gatttqata  | ctgttatgaa  | 420  |
| 200 | ttgtaataat  | aataattttg  | aagaagagg    | tttgcagg    | gtttgagaac | tgctgttcta  | 480  |
| 202 | gccccacgtg  | gtgggtttt   | cgtcattttg   | ggtttttatg  | aggcagagtc | ttatgttagcc | 540  |
| 204 | caggctagca  | gcotagaatg  | tgctacttgc   | ctgaggaa    | accttggaa  | ttctgaggac  | 600  |
| 206 | tggagagact  | ggcttaqtcc  | tcaagaaact   | ggaaataatgt | ggagttttgc | tacttgggg   | 660  |
| 208 | ttcccttttc  | tcaaaaccc   | tttctactctt  | tttccaccc   | gtcgcccccc | taacactaaa  | 720  |
| 210 | taagaagag   | aaaggggago  | atagaggggaa  | aaagaaaccc  | ctgaaaaacg | tcagtagttg  | 780  |
| 212 | gcaaggggg   | gtgacataatg | ttgtcatttag  | accacatctt  | ggttggaa   | gggagtcaag  | 840  |
| 214 | ttccttgggg  | caagtttgc   | ctttgtgtt    | acgatatcta  | attttttctc | cctgttgctt  | 900  |
| 216 | cgttttgcg   | aacaacgact  | tgataaccca   | caatggacca  | tcaaccaacc | aaccaaccat  | 960  |
| 218 | <210>       | SEQ ID NO:  | 6            |             |            |             |      |
| 219 | <211>       | LENGTH:     | 1090         | base pairs  |            |             |      |
| 220 | <212>       | TYPE:       | DNA          |             |            |             |      |
| 221 | <213>       | ORGANISM:   | Homo sapiens |             |            |             |      |
| 223 | <400>       | SEQUENCE:   | 6            |             |            |             |      |
| 224 | ttgtctctgg  | tgttactgt   | tttccat      | ctgacagtgg  | tttgcaccc  | tatacgcc    | 60   |
| 226 | tgtgtcgg    | gtgtgtaga   | cctat        | tttcc       | tttcc      | cagccat     | 120  |
| 228 | ttgtctact   | gtcagatgt   | tgactgttcc   | tttcc       | tttcc      | gtctctgt    | 180  |
| 230 | gcaggaacca  | gaaggccctg  | ttcc         | tttcc       | tttcc      | ctacgcac    | 240  |
| 232 | tgtgtctgg   | ttgttcc     | ttc          | tttcc       | tttcc      | tttcc       | 300  |
| 234 | ttcttagtat  | gtcttcc     | ttc          | tttcc       | tttcc      | tttcc       | 360  |
| 236 | ggagcttt    | gaccagg     | tttcc        | tttcc       | tttcc      | tttcc       | 420  |
| 238 | tttgctgt    | gtcaat      | tttcc        | tttcc       | tttcc      | tttcc       | 480  |
| 240 | ttggcaag    | gtggca      | tttcc        | tttcc       | tttcc      | tttcc       | 540  |
| 242 | acttcgtggc  | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 600  |
| 244 | atcaggcaa   | ggtttgg     | tttcc        | tttcc       | tttcc      | tttcc       | 660  |
| 246 | ttgccttgc   | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 720  |
| 248 | ccctcgg     | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 780  |
| 250 | gtatgtttt   | aaaatctaa   | tttcc        | tttcc       | tttcc      | tttcc       | 840  |
| 252 | ttgttcc     | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 900  |
| 254 | atccaaatc   | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 960  |
| 256 | ctcgcc      | tttcc       | tttcc        | tttcc       | tttcc      | tttcc       | 1020 |
| 258 | tgaatctaa   | ggagaag     | tttcc        | tttcc       | tttcc      | tttcc       | 1080 |

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260 tgcatccaca 1090

VERIFICATION SUMMARY  
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Input Set : A:\Sequence\_Listing.txt  
Output Set: N:\CRF3\10302000\I101423A.raw

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L:6 M:283 W: Missing Blank Line separator, <130> field identifier  
L:7 M:283 W: Missing Blank Line separator, <140> field identifier  
L:11 M:283 W: Missing Blank Line separator, <160> field identifier  
L:59 M:283 W: Missing Blank Line separator, <400> field identifier